

# SEQUENCE LISTING

<110> Stephen M. Allen  
Saverio C. Falco  
Catherine J. Thorpe

<120> Genes Encoding Sulfate Assimilation Proteins

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<150> 60/092,833

<151> 14-07-1998

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<212> DNA

<213> Zea mays

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Lys Val Ala Pro Pro Pro Ala Arg Ser Thr Ala Ser Lys Met Lys Val  
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Gly Gln Pro Pro Gly Thr Gln Trp Leu Met Ala Val Arg Tyr Leu Phe  
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Pro Ile Leu Asp Trp Val Pro Ser Tyr Ser Leu Ser Leu Phe Lys Ser  
115 120 125

Asp Leu Val Ala Gly Leu Thr Ile Ala Ser Leu Ala Ile Pro Gln Gly  
130 135 140

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Ser Ser Phe Val Pro Pro Met Val Tyr Ala Val Leu Gly Ser Ser Arg  
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Gln Leu Ala Phe Thr Ser Thr Leu Phe Ala Gly Leu Val Gln Ala Ser  
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Phe Leu Ile Asp Ile Lys Lys Ser Ile Glu Lys Arg Gly Leu Glu Leu  
625 630 635 640

Val Leu Val Asn Pro Thr Gly Glu Val Met Glu Lys Ile Gln Arg Ala  
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<212> DNA  
<213> Zea mays

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 <212> PRT  
 <213> Zea mays

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 Gly Ser Ser Lys Asp Leu Ala Val Gly Thr Val Ala Val Ala Ser Leu  
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 Phe Gln Ala Ser Leu Gly Leu Leu Arg Leu Gly Phe Ile Val Asp Leu  
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 Leu Ser His Ala Thr Ile Val Gly Phe Met Ala Gly Ala Ala Thr Val  
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 Val Cys Leu Gln Gln Leu Lys Gly Met Leu Gly Leu Val His Phe Thr  
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 Thr Ser Thr Asp Val Val Ser Val Met Glu Ser Val Phe Ser Gln Thr  
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 165 170 175  
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 195 200 205  
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 Tyr Leu Lys Lys Gly Leu Asn Pro Pro Ser Val Thr Ser Leu Gln Phe  
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 245 250 255  
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Met Lys Leu Leu Tyr Tyr Thr Pro Met Ala Val Leu Ala Ser Ile Ile  
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Leu Ser Ala Leu Pro Gly Leu Ile Asp Ile Lys Glu Ala Cys Ser Ile  
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Trp Lys Ile Asp Lys Met Asp Phe Leu Thr Cys Leu Gly Ala Phe Val  
115 120 125  
Gly Val Leu Phe Gly Ser Val Glu Ile Gly Leu Ala Val Ala Leu Gly  
130 135 140

Ile Ser Phe Ala Lys Ile Ile Ile Gln Ser Leu Arg Pro Gln Val Glu  
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 180 185 190  
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 Glu Arg Ile Gln Ala Val Val Leu Asp Met Ser Ser Val Val Asn Ile  
 225 230 235 240  
 Asp Thr Ser Gly Leu Thr Ala Leu Glu Glu Ile His Lys Glu Leu Val  
 245 250 255  
 Ser Leu Gly Leu Gln Met Ala Ile Ala Ser Pro Gly Trp Lys Ala Val  
 260 265 270  
 Gln Lys Met Lys Val Ser Gln Val Val Asp Arg Val Gly Gln Asp Trp  
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 Leu Leu Phe Val Ser Arg Pro Arg Thr Ser Thr Leu Gly Leu Ile Pro  
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 Asp Ser Thr Ile Tyr Arg Ser Met Asp Gln Tyr Gln Asn Ala Lys Ser  
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 Val Pro Gly Ile Leu Ile Leu Gln Ile Glu Ala Pro Ile Tyr Phe Ala  
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 Asn Ser Ser Tyr Leu Arg Glu Arg Ile Val Arg Trp Val Asp Glu Glu  
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 Glu Asp Arg Leu Lys Ser Leu Lys Glu Asn Asp Leu Gln Tyr Val Ile  
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 Leu Gly Glu Val Lys Lys Val Met Glu Arg Arg Gly Leu Lys Leu Val  
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 180 185 190  
 Ala Val Gly Ala Cys Asn Phe Met Leu His Thr Tyr Lys Asn Ala Glu  
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tatcaattag attggaacaa ggagatggaa cacttggaac tatgaacata caaggtcaat 480
gana 484

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<210> 10
<211> 147
<212> PRT
<213> Oryza sativa

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<220>
<221> UNSURE
<222> (17)
<223> Xaa = any amino acid

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<220>
<221> UNSURE
<222> (131)
<223> Xaa = any amino acid

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 1          5          10          15

Xaa Ser Pro Asn Arg Ala His Asn Arg Trp Asn Trp Gln Thr Ile Val
      20          25          30

Ile Gly Ile Thr Phe Leu Ala Phe Leu Leu Leu Ala Lys Tyr Ile Gly
 35          40          45

Lys Lys Asn Arg Lys Phe Phe Trp Val Pro Ala Ile Ala Pro Ile Thr
 50          55          60

Ser Val Ile Leu Ala Thr Leu Phe Val Phe Ile Thr Arg Ala Asp Lys
 65          70          75          80

Gln Gly Val Gln Ile Val Asn His Ile Lys Lys Gly Ile Asn Pro Ser
      85          90          95

Ser Val His Lys Ile Tyr Phe Thr Gly Pro Phe Val Ala Lys Gly Phe
      100          105          110

Lys Ile Gly Val Ile Ser Ala Met Ile Gly Leu Thr Glu Ala Val Ala
      115          120          125

Ile Gly Xaa Thr Phe Ala Ala Leu Lys Asp Tyr Gln Leu Asp Trp Asn
      130          135          140

Lys Glu Met
145

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<210> 11
<211> 510

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<212> DNA  
 <213> Oryza sativa

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 acggcggcgg aggaggggcg ggggcggcga gggtgccgat gccggcggcg aagccgttcc 180  
 tggagacgct ggggggggaa atgaaggaga cattcctgcc ggacgacccg ttcaggggtgg 240  
 tgcggcgggga gcgcgggtgc gggcggcgcg cggcggcggc gctccggtac gtgttcccgt 300  
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 caccattgcc agcctcagca tcccgcaagg gatcagctag ccaagctcgc aactccctcg 420  
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<210> 12  
 <211> 98  
 <212> PRT  
 <213> Oryza sativa

<400> 12  
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 20 25 30  
 Gly Asn Met Lys Glu Thr Phe Leu Pro Asp Asp Pro Phe Arg Val Val  
 35 40 45  
 Arg Arg Glu Arg Gly Cys Gly Arg Arg Ala Ala Ala Ala Leu Arg Tyr  
 50 55 60  
 Val Phe Pro Phe Met Glu Trp Ala Pro Ser Tyr Thr Leu Gly Thr Leu  
 65 70 75 80  
 Lys Ser Asp Leu Ile Ala Gly Thr Pro Leu Pro Ala Ser Ala Ser Arg  
 85 90 95  
 Lys Gly

<210> 13  
 <211> 493  
 <212> DNA  
 <213> Oryza sativa

<400> 13  
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 gcctctgctg tcgggtcatcc tgtccaccgc tgcggtctac gcgacaaggg ctgacaggca 180  
 cggcgtcaag atcatccaga aggtgcacgc gggcctaaac ccaagctccg tgggaagcaga 240  
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 tcgccctcac ggaagctatc gccgttgccc gatctttcgc ctccgtaaga gggtagagac 360  
 tcgacggcaa caaggagatg ctggccatgg ggttctccaa cgttgctggt tctctgtcct 420  
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<210> 14  
 <211> 159  
 <212> PRT

<213> Oryza sativa

<220>

<221> UNSURE

<222> (74)

<223> Xaa = any amino acid

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Trp His Pro Gly Asn Phe Leu Ile Gly Cys Ser Phe Leu Ile Phe Ile  
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Leu Thr Thr Arg Phe Ile Gly Arg Arg Tyr Lys Lys Leu Phe Trp Leu  
20 25 30

Ser Ala Ile Ser Pro Leu Leu Ser Val Ile Leu Ser Thr Ala Ala Val  
35 40 45

Tyr Ala Thr Arg Ala Asp Arg His Gly Val Lys Ile Ile Gln Lys Val  
50 55 60

His Ala Gly Leu Asn Pro Ser Ser Val Xaa Gln Ile His Leu Asn Gly  
65 70 75 80

Pro His Thr Thr Glu Cys Ala Gln Asp Arg Arg His Leu Arg Ile Ile  
85 90 95

Ala Leu Thr Glu Ala Ile Ala Val Gly Arg Ser Phe Ala Ser Val Arg  
100 105 110

Gly Tyr Arg Leu Asp Gly Asn Lys Glu Met Leu Ala Met Gly Phe Ser  
115 120 125

Asn Val Ala Gly Ser Leu Ser Ser Cys Tyr Val Ala Thr Gly Ser Phe  
130 135 140

Ser Arg Thr Ala Val Asn Phe Ser Gly Gly Gly Gln Ser Thr Val  
145 150 155

<210> 15

<211> 2067

<212> DNA

<213> Glycine max

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acttcagtat	ctcttcccca	ttttcgactg	ggccccaac	tacaatctta	cccttctccg	180
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gatatactcg	ctgcttgga	gttctagaca	tcttggtggt	ggacctgttt	ccattgcgtc	360
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attaaggcta	ggcttcgtaa	ttgattttct	gtcgaaggca	acgctggttg	gattcacagg	540
cgggtgctgcc	attattgtgt	cactgcagca	gctgaaaggt	ttacttgga	tagtgcactt	600
taccagcaag	atgcaaataa	ttccagtaac	gatctctgtt	ttcaagcaaa	gacacgagt	660
gtcatggcaa	accattcttt	tgggattcgg	cttcctggtc	ttcttgctga	caacaaggca	720
cattagtttg	aggaaaccaa	aactattctg	ggtttcagca	gctgccccat	tgacatcagt	780
tattctgtca	accattttag	tctttcttct	gagaaataag	actcatcaaa	tttcagttat	840
tgggcactta	caaaggagg	ttaatccacc	atcagcaaac	atgttatact	tcaatgggtcc	900
ttacttgggt	cttgctatca	aaactggcat	catcacaggg	atcttatctc	tcactgaagg	960

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aattgcagta gggagaacat ttgcttcact taagaactac caggtggatg gaaacaaaga 1020
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gggatccttt tctcgatcgg ctgttaacta taatgctgga gcacagacaa cagtttcaaa 1140
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gccttgattt cccatcaatg ttgttcaagg acttatatat ggggataaac tctctaacct 1920
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aaaaaaaaa aaaaaaaaaa aaaaaaa 2067

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<210> 16
<211> 621
<212> PRT
<213> Glycine max

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His Glu Pro His Gln Thr Thr Leu His Lys Leu Arg His Arg Val Ser
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          20          25          30

Arg Phe Lys Lys Phe Leu Leu Ala Leu Gln Tyr Leu Phe Pro Ile Phe
          35          40          45

Asp Trp Ala Pro Asn Tyr Asn Leu Thr Leu Leu Arg Ser Asp Leu Ile
          50          55          60

Ser Gly Leu Thr Ile Ala Ser Leu Ala Ile Pro Gln Gly Ile Ser Tyr
          65          70          75          80

Ala Lys Leu Ala Asn Leu Pro Pro Ile Leu Gly Leu Tyr Ser Ser Phe
          85          90          95

Val Pro Pro Leu Ile Tyr Ser Leu Leu Gly Ser Ser Arg His Leu Gly
          100          105          110

Val Gly Pro Val Ser Ile Ala Ser Leu Val Met Gly Ser Met Leu Ser
          115          120          125

Asp Lys Ile Ser Tyr Thr Gln Glu Pro Ile Leu Tyr Leu Gly Leu Ala
          130          135          140

Phe Thr Ala Thr Phe Phe Ala Gly Val Phe Gln Ala Ser Leu Gly Ile
          145          150          155          160

Leu Arg Leu Gly Phe Val Ile Asp Phe Leu Ser Lys Ala Thr Leu Val
          165          170          175

Gly Phe Thr Gly Gly Ala Ala Ile Ile Val Ser Leu Gln Gln Leu Lys

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180					185					190					
Gly	Leu	Leu	Gly	Ile	Val	His	Phe	Thr	Ser	Lys	Met	Gln	Ile	Ile	Pro
	195						200					205			
Val	Thr	Ile	Ser	Val	Phe	Lys	Gln	Arg	His	Glu	Trp	Ser	Trp	Gln	Thr
	210					215					220				
Ile	Leu	Leu	Gly	Phe	Gly	Phe	Leu	Val	Phe	Leu	Leu	Thr	Thr	Arg	His
225					230					235					240
Ile	Ser	Leu	Arg	Lys	Pro	Lys	Leu	Phe	Trp	Val	Ser	Ala	Ala	Ala	Pro
				245					250					255	
Leu	Thr	Ser	Val	Ile	Leu	Ser	Thr	Ile	Leu	Val	Phe	Leu	Leu	Arg	Asn
			260					265					270		
Lys	Thr	His	Gln	Ile	Ser	Val	Ile	Gly	His	Leu	Pro	Lys	Gly	Val	Asn
		275					280					285			
Pro	Pro	Ser	Ala	Asn	Met	Leu	Tyr	Phe	Asn	Gly	Pro	Tyr	Leu	Gly	Leu
	290					295					300				
Ala	Ile	Lys	Thr	Gly	Ile	Ile	Thr	Gly	Ile	Leu	Ser	Leu	Thr	Glu	Gly
305					310					315					320
Ile	Ala	Val	Gly	Arg	Thr	Phe	Ala	Ser	Leu	Lys	Asn	Tyr	Gln	Val	Asp
				325					330					335	
Gly	Asn	Lys	Glu	Met	Met	Ala	Ile	Gly	Leu	Met	Asn	Ile	Ala	Gly	Ser
			340					345					350		
Cys	Ser	Ser	Cys	Tyr	Val	Thr	Thr	Gly	Ser	Phe	Ser	Arg	Ser	Ala	Val
		355					360					365			
Asn	Tyr	Asn	Ala	Gly	Ala	Gln	Thr	Thr	Val	Ser	Asn	Ile	Ile	Met	Ala
	370					375					380				
Ala	Ala	Val	Leu	Val	Thr	Leu	Leu	Phe	Leu	Met	Pro	Leu	Phe	Tyr	Tyr
385					390					395					400
Thr	Pro	Asn	Val	Val	Leu	Ala	Ala	Ile	Ile	Ile	Thr	Ala	Val	Ile	Gly
				405					410					415	
Leu	Ile	Asp	Tyr	Gln	Ser	Ala	Tyr	Lys	Leu	Trp	Lys	Val	Asp	Lys	Leu
		420						425					430		
Asp	Phe	Leu	Ala	Cys	Leu	Cys	Ser	Phe	Phe	Gly	Val	Leu	Phe	Ile	Ser
		435					440					445			
Val	Pro	Leu	Gly	Leu	Gly	Ile	Ala	Val	Ile	Ile	Ser	Val	Leu	Lys	Ile
	450					455					460				
Leu	Leu	His	Val	Thr	Arg	Pro	Asn	Thr	Leu	Val	Leu	Gly	Asn	Ile	Pro
465					470					475					480
Gly	Thr	Gln	Ile	Phe	His	Asn	Ile	Asn	Gln	Tyr	Lys	Lys	Ala	Leu	Arg
				485					490					495	
Val	Pro	Ser	Phe	Leu	Ile	Leu	Ala	Val	Glu	Ser	Pro	Ile	Tyr	Phe	Ala

500	505	510
Asn Ser Thr Tyr Leu Gln Glu Arg Ile Leu Arg Trp Val Arg Glu Glu		
515	520	525
Glu Glu His Ile Lys Ala Asn Asn Gly Ala Pro Leu Lys Cys Ile Ile		
530	535	540
Leu Asp Met Thr Ala Val Thr Ala Thr Asp Thr Ser Gly Leu Asp Thr		
545	550	555
Leu Cys Glu Leu Arg Lys Met Leu Glu Lys Arg Ser Leu Glu Phe Val		
565	570	575
Leu Ala Asn Pro Val Gly Asn Val Met Glu Lys Leu His Lys Ser Asn		
580	585	590
Ile Leu Asp Ser Phe Gly Leu Lys Gly Val Tyr Leu Thr Val Gly Glu		
595	600	605
Ala Val Thr Asp Ile Ser Ser Ile Trp Lys Ala Gln Pro		
610	615	620

<210> 17  
 <211> 2449  
 <212> DNA  
 <213> Glycine max

<400> 17

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agtgcaccaa	gtcagaggttc	caccgccaca	gccgtttttc	aagtctctaa	agtactcttt	180
gaaggagact	ttcttccttg	atgacccttt	gaggcagttc	aagaacaagc	cagcttccaa	240
gaagttcatg	cttggcccttc	agttcttctt	ccccattttc	gaatgggctc	ccaaatacac	300
ctttcagttc	ttgaaagctg	acctcatagc	tggcatcacc	atcgctagct	tggccattcc	360
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ctttatacca	ccattgattt	atgcatgat	gggtagctcg	agggatttgg	cagtggggac	480
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tgttttggtc aatcctgtaa gtgaagtgat gaagaaactg aacaaatcga agttccaaaa 1920
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caatctacgt gcaagcaaaa cgaaccctaaa gaaagatgaa acagaggggtt ggaacaatgt 2040
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<210> 18
<211> 680
<212> PRT
<213> Glycine max

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Arg Asn Thr Ile Ile Glu Asp Met Gly Ser Val Asp Tyr Glu Tyr Pro
          20          25          30

Leu Gly Met Asn Asn Phe Glu Arg Val His Gln Val Glu Val Pro Pro
          35          40          45

Pro Gln Pro Phe Phe Lys Ser Leu Lys Tyr Ser Leu Lys Glu Thr Phe
          50          55          60

Phe Pro Asp Asp Pro Leu Arg Gln Phe Lys Asn Lys Pro Ala Ser Lys
          65          70          75          80

Lys Phe Met Leu Gly Leu Gln Phe Phe Phe Pro Ile Phe Glu Trp Ala
          85          90          95

Pro Lys Tyr Thr Phe Gln Phe Leu Lys Ala Asp Leu Ile Ala Gly Ile
          100          105          110

Thr Ile Ala Ser Leu Ala Ile Pro Gln Gly Ile Ser Tyr Ala Lys Leu
          115          120          125

Ala Asn Leu Pro Pro Ile Leu Gly Leu Tyr Ser Ser Phe Ile Pro Pro
          130          135          140

Leu Ile Tyr Ala Met Met Gly Ser Ser Arg Asp Leu Ala Val Gly Thr
          145          150          155          160

Val Ala Val Gly Ser Leu Leu Met Gly Ser Met Leu Ser Asn Ala Val
          165          170          175

Asp Pro Asn Glu Asp Pro Lys Leu Tyr Leu His Leu Ala Phe Thr Ala
          180          185          190

Thr Leu Phe Ala Gly Val Phe Gln Ala Ala Leu Gly Leu Phe Arg Leu
          195          200          205

Gly Leu Ile Val Asp Phe Leu Ser His Ala Thr Ile Ile Gly Phe Met
          210          215          220

Gly Gly Ala Ala Thr Val Val Cys Leu Gln Gln Leu Lys Ser Ile Leu

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225					230					235					240
Gly	Leu	Glu	His	Phe	Thr	His	Gly	Ala	Asp	Ile	Ile	Ser	Val	Met	Arg
				245					250					255	
Ser	Val	Phe	Thr	Gln	Thr	His	Glu	Trp	Arg	Trp	Glu	Ser	Ala	Val	Leu
			260					265					270		
Gly	Cys	Val	Phe	Ile	Phe	Phe	Leu	Leu	Ser	Thr	Arg	Tyr	Phe	Ser	Lys
		275					280					285			
Lys	Arg	Pro	Arg	Phe	Phe	Trp	Val	Ser	Ala	Met	Ala	Pro	Leu	Thr	Ser
	290					295					300				
Val	Ile	Leu	Gly	Ser	Leu	Leu	Val	Tyr	Phe	Thr	His	Ala	Glu	Lys	His
305					310					315					320
Gly	Val	Glu	Val	Ile	Gly	Glu	Leu	Lys	Lys	Gly	Leu	Asn	Pro	Pro	Ser
				325					330					335	
Leu	Thr	Asn	Leu	Val	Phe	Val	Ser	Pro	Tyr	Met	Thr	Thr	Ala	Val	Lys
			340					345					350		
Thr	Gly	Ile	Val	Val	Gly	Ile	Ile	Ser	Leu	Ala	Glu	Gly	Ile	Ala	Val
		355					360					365			
Gly	Arg	Ser	Phe	Ala	Met	Tyr	Lys	Asn	Tyr	Asn	Ile	Asp	Gly	Asn	Lys
	370					375					380				
Glu	Met	Ile	Ala	Ile	Gly	Thr	Met	Asn	Val	Val	Gly	Ser	Phe	Thr	Ser
385					390					395					400
Cys	Tyr	Leu	Thr	Thr	Gly	Pro	Phe	Ser	Arg	Ser	Ala	Val	Asn	Tyr	Asn
				405					410					415	
Ala	Gly	Cys	Lys	Thr	Ala	Ala	Ser	Asn	Ile	Ile	Met	Ser	Leu	Ala	Val
			420					425					430		
Met	Leu	Thr	Leu	Leu	Phe	Leu	Thr	Pro	Leu	Phe	His	Tyr	Thr	Pro	Leu
		435					440					445			
Val	Val	Leu	Ser	Ala	Ile	Ile	Val	Ser	Ala	Met	Leu	Gly	Leu	Ile	Asp
	450					455					460				
Tyr	Glu	Ala	Ala	Ile	His	Leu	Phe	Lys	Val	Asp	Lys	Phe	Asp	Phe	Val
465					470					475					480
Val	Cys	Met	Ser	Ala	Tyr	Ile	Gly	Val	Val	Phe	Gly	Ser	Val	Glu	Ile
				485					490					495	
Gly	Leu	Val	Ile	Ala	Ile	Val	Ile	Ser	Val	Leu	Arg	Val	Leu	Leu	Phe
			500					505					510		
Ile	Ala	Arg	Pro	Arg	Thr	Phe	Val	Leu	Gly	Asn	Ile	Pro	Asn	Ser	Val
		515					520					525			
Ile	Tyr	Arg	Asn	Val	Glu	His	Tyr	Gln	Asn	Ala	Lys	His	Val	Pro	Gly
	530					535					540				
Met	Leu	Ile	Leu	Glu	Ile	Asp	Ala	Pro	Ile	Tyr	Phe	Ala	Asn	Ala	Ser

545                      550                      555                      560  
 Tyr Leu Arg Glu Arg Ile Thr Arg Trp Ile Asp Glu Glu Glu Glu Arg  
                                  565                      570                      575  
 Ile Lys Ala Thr Gly Glu Thr Ser Leu Gln Tyr Val Ile Ile Asp Met  
                                  580                      585                      590  
 Ser Ala Val Gly Asn Ile Asp Thr Ser Gly Ile Ser Met Leu Glu Glu  
                                  595                      600                      605  
 Val Lys Lys Ile Thr Glu Arg Arg Glu Leu Gln Leu Val Leu Val Asn  
                                  610                      615                      620  
 Pro Val Ser Glu Val Met Lys Lys Leu Asn Lys Ser Lys Phe Gln Asn  
 625                                   630                                   635                                   640  
 His Leu Gly Lys Lys Trp Ile Tyr Leu Thr Val Glu Glu Ala Val Gly  
                                  645                                   650                                   655  
 Ala Cys Asn Phe Asn Leu Arg Ala Ser Lys Thr Asn Pro Lys Lys Asp  
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 Glu Thr Glu Gly Trp Asn Asn Val  
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 <211> 2311  
 <212> DNA  
 <213> Triticum aestivum

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 gaggcagcag atgaacctag catcaccaca cagacacccc ccaatgaccc atctcaagca 180  
 ccgctggtgt acaaagtggg ctatccccct ccgaagaact tggccacaga gtttacagaa 240  
 acattgaggg agacttttctt ccacgacaac ccgctgcgtc agtataaggg ccaatccgga 300  
 ccgaggaggt tcatgatggg gctggagttc ttgtttccta tatttgggtg gggtagggat 360  
 tacagtctca acaagttcaa aggcgatctg attgccggat tgaccatcgc aagtctctgt 420  
 attcctcagg acattggcta ttcgaagctt gctaattctg atccgcagta tgggctttac 480  
 tccagcttca ttctctcatt gatctatgct gcaatgggta gctcaaggga tatagcgatt 540  
 ggtccagttg ctgtgggttc tcttttgata gggtcacttc tacaagctga ggttgaccat 600  
 gtcaaaaaca aggaggaata catgcgcctc gctttcacgg caaccttctt cgctgggtatc 660  
 actcaagcag ccttaggatt tctaaggtta ggattcctta tagagttctt gtcgcatgct 720  
 gcgattgtcg gattcatggg gggagctgcc attactattg ccctgcagca gctgaaatac 780  
 gtggtgggca tcgcaaactt tacaaggaaa accgacatag tttctgtcat ggaatctgtc 840  
 tggagatcag ttcatacagg gtggaactgg cagacaattg tgattggcgt atctttcctg 900  
 gttttccttc tgtttgcgaa gtacatcgga aagaagaaaa ggaagctttt ctgggtgcc 960  
 gctattgtct ctataatttc agtgattcta gcaacatttt ttgtatacat tactcgtgcc 1020  
 gacaagcaag gagttcagat agtgaagcac attgaacagg gaatcaaccc atcatcagta 1080  
 cacaagattt atttcaccgg cccatttggt gcaaaagggt tcaagatcgg tgttggttgc 1140  
 ggcatagttg gtttgacaga agctgtagct attggaagga catttgctgc tatgaaggac 1200  
 taccagttag atggaaacaa ggagatggta gcacttgga ccatgaacat agtaggctca 1260  
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 gcggtgatcg gccttggtga ctacgaagca gcaattctca tctggaaagt tgacaaattg 1500  
 gacttcattg cttgcatggg agcttttttc ggtgttgttt ttgtatccgt tgagattggc 1560  
 ctcttgattg ctgtagcaat ctcatgtgcc aaaatacttc ttcaagtaac aaggccaagg 1620  
 acagccctac ttggaaacct tcccggcacc actatatacc ggaacatcag ccagtatcca 1680

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gaagcaaaac ttactcctgg ggtggtgatt gtgagggttg attctgctat ttatttttcc 1740
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ctcaccgagc acattggaag cagcaatata ttcctcgcg tctctgacgc tgtgcgattc 2040
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gagaatagtt tggccgctcc ctgtgatcta agctgggaca gcgcaatatg atgtggcttt 2160
gtggccaatg tagaaacata taataagtta aggcaatcac cggagcttct ccggtttact 2220
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aaaaaaaaa aaaaaaaaaa aaaaaaaaaa a a 2311

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<212> PRT
<213> Triticum aestivum

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Thr Gln Thr Pro Pro Asn Asp Pro Ser Gln Ala Pro Leu Val Tyr Lys
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Val Gly Tyr Pro Pro Pro Lys Asn Leu Ala Thr Glu Phe Thr Glu Thr
          35          40          45

Leu Arg Glu Thr Phe Phe His Asp Asn Pro Leu Arg Gln Tyr Lys Gly
          50          55          60

Gln Ser Gly Pro Arg Arg Phe Met Met Gly Leu Glu Phe Leu Phe Pro
          65          70          75          80

Ile Phe Gly Trp Gly Arg Asp Tyr Ser Leu Asn Lys Phe Lys Gly Asp
          85          90          95

Leu Ile Ala Gly Leu Thr Ile Ala Ser Leu Cys Ile Pro Gln Asp Ile
          100          105          110

Gly Tyr Ser Lys Leu Ala Asn Leu Asp Pro Gln Tyr Gly Leu Tyr Ser
          115          120          125

Ser Phe Ile Pro Pro Leu Ile Tyr Ala Ala Met Gly Ser Ser Arg Asp
          130          135          140

Ile Ala Ile Gly Pro Val Ala Val Val Ser Leu Leu Ile Gly Ser Leu
          145          150          155          160

Leu Gln Ala Glu Val Asp His Val Lys Asn Lys Glu Glu Tyr Met Arg
          165          170          175

Leu Ala Phe Thr Ala Thr Phe Phe Ala Gly Ile Thr Gln Ala Ala Leu
          180          185          190

Gly Phe Leu Arg Leu Gly Phe Leu Ile Glu Phe Leu Ser His Ala Ala
          195          200          205

Ile Val Gly Phe Met Gly Gly Ala Ala Ile Thr Ile Ala Leu Gln Gln
          210          215          220

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Leu	Lys	Tyr	Val	Leu	Gly	Ile	Ala	Asn	Phe	Thr	Arg	Lys	Thr	Asp	Ile	225	230	235	240
Val	Ser	Val	Met	Glu	Ser	Val	Trp	Arg	Ser	Val	His	His	Gly	Trp	Asn	245	250	255	
Trp	Gln	Thr	Ile	Val	Ile	Gly	Val	Ser	Phe	Leu	Val	Phe	Leu	Leu	Phe	260	265	270	
Ala	Lys	Tyr	Ile	Gly	Lys	Lys	Lys	Arg	Lys	Leu	Phe	Trp	Val	Pro	Ala	275	280	285	
Ile	Ala	Pro	Ile	Ile	Ser	Val	Ile	Leu	Ala	Thr	Phe	Phe	Val	Tyr	Ile	290	295	300	
Thr	Arg	Ala	Asp	Lys	Gln	Gly	Val	Gln	Ile	Val	Lys	His	Ile	Glu	Gln	305	310	315	320
Gly	Ile	Asn	Pro	Ser	Ser	Val	His	Lys	Ile	Tyr	Phe	Thr	Gly	Pro	Phe	325	330	335	
Val	Ala	Lys	Gly	Phe	Lys	Ile	Gly	Val	Val	Cys	Gly	Ile	Val	Gly	Leu	340	345	350	
Thr	Glu	Ala	Val	Ala	Ile	Gly	Arg	Thr	Phe	Ala	Ala	Met	Lys	Asp	Tyr	355	360	365	
Gln	Leu	Asp	Gly	Asn	Lys	Glu	Met	Val	Ala	Leu	Gly	Thr	Met	Asn	Ile	370	375	380	
Val	Gly	Ser	Met	Thr	Ser	Cys	Tyr	Val	Thr	Thr	Gly	Ser	Phe	Ser	Arg	385	390	395	400
Ser	Ala	Val	Asn	Phe	Met	Ala	Gly	Cys	Lys	Thr	Pro	Val	Ser	Asn	Val	405	410	415	
Val	Met	Ser	Val	Val	Val	Leu	Leu	Thr	Leu	Leu	Val	Ile	Thr	Pro	Leu	420	425	430	
Phe	Lys	Tyr	Thr	Pro	Asn	Ala	Ile	Leu	Gly	Ser	Ile	Ile	Ile	Ser	Ala	435	440	445	
Val	Ile	Gly	Leu	Val	Asp	Tyr	Glu	Ala	Ala	Ile	Leu	Ile	Trp	Lys	Val	450	455	460	
Asp	Lys	Leu	Asp	Phe	Ile	Ala	Cys	Met	Gly	Ala	Phe	Phe	Gly	Val	Val	465	470	475	480
Phe	Val	Ser	Val	Glu	Ile	Gly	Leu	Leu	Ile	Ala	Val	Ala	Ile	Ser	Phe	485	490	495	
Ala	Lys	Ile	Leu	Leu	Gln	Val	Thr	Arg	Pro	Arg	Thr	Ala	Leu	Leu	Gly	500	505	510	
Asn	Leu	Pro	Gly	Thr	Thr	Ile	Tyr	Arg	Asn	Ile	Ser	Gln	Tyr	Pro	Glu	515	520	525	
Ala	Lys	Leu	Thr	Pro	Gly	Val	Val	Ile	Val	Arg	Val	Asp	Ser	Ala	Ile	530	535	540	

Tyr Phe Ser Asn Ser Asn Tyr Val Arg Glu Arg Ile Leu Arg Trp Leu  
 545 550 555 560  
 Thr Asp Glu Glu Asp Arg Ala Lys Ala Val Gly Leu Pro Lys Ile Ser  
 565 570 575  
 Phe Leu Ile Val Glu Met Ser Pro Val Ile Asp Ile Asp Thr Ser Gly  
 580 585 590  
 Ile His Ala Leu Glu Asp Leu Tyr Lys Asn Leu Gln Lys Lys Asp Met  
 595 600 605  
 Gln Leu Ile Leu Ser Asn Pro Gly Ser Val Val Ile Glu Lys Leu Gln  
 610 615 620  
 Ala Ser Lys Leu Thr Glu His Ile Gly Ser Ser Asn Ile Phe Leu Ala  
 625 630 635 640  
 Val Ser Asp Ala Val Arg Phe Cys Thr Thr Lys Ser Met Gln Glu Pro  
 645 650 655

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 <211> 2022  
 <212> DNA  
 <213> Triticum aestivum

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 tcgccgcccg catcactgtc ggcgtcatgc ttgtgcctca ggcaatgtca tatgcaaagc 180  
 tggctgggct tcacccaatt tatgggctct acacaggctt tgtccacta tttgtctacg 240  
 cgatttttgg gtcctcacga caattagcag taggtccagt ggcacttgct tctctgctag 300  
 tgtccaatgt tcttgggggt atagttaatt catctagtga gctgtacacg gaattagcca 360  
 tattattggc attcatgggt ggaatactgg aatgcttgat ggcattgcta agacttggct 420  
 ggcttattcg ttctattagc cattctgtaa tatctggatt cactacagct tcggccatcg 480  
 taattgggtt gtcccaaata aagtatttct tgggttacag tgttacaaga agtagcaaaa 540  
 ttataccact tattgagagt ataattgctg gaatagatca gttctcctgg cctccatttg 600  
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 aaaaattacg ttctctgaga gcttctggct cactaacagc tgttggttctt ggaacattgt 720  
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2022

<210> 22

<211> 593

<212> PRT

<213> Triticum aestivum

<400> 22

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Ala Glu Ala Ala Leu Pro Cys Leu Ala Trp Met Arg Ser Tyr Arg Trp  
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Lys Glu Asp Phe Gln Ala Asp Leu Ala Ala Gly Ile Thr Val Gly Val  
35 40 45

Met Leu Val Pro Gln Ala Met Ser Tyr Ala Lys Leu Ala Gly Leu His  
50 55 60

Pro Ile Tyr Gly Leu Tyr Thr Gly Phe Val Pro Leu Phe Val Tyr Ala  
65 70 75 80

Ile Phe Gly Ser Ser Arg Gln Leu Ala Val Gly Pro Val Ala Leu Val  
85 90 95

Ser Leu Leu Val Ser Asn Val Leu Gly Gly Ile Val Asn Ser Ser Ser  
100 105 110

Glu Leu Tyr Thr Glu Leu Ala Ile Leu Leu Ala Phe Met Val Gly Ile  
115 120 125

Leu Glu Cys Leu Met Ala Leu Leu Arg Leu Gly Trp Leu Ile Arg Phe  
130 135 140

Ile Ser His Ser Val Ile Ser Gly Phe Thr Thr Ala Ser Ala Ile Val  
145 150 155 160

Ile Gly Leu Ser Gln Ile Lys Tyr Phe Leu Gly Tyr Ser Val Thr Arg  
165 170 175

Ser Ser Lys Ile Ile Pro Leu Ile Glu Ser Ile Ile Ala Gly Ile Asp  
180 185 190

Gln Phe Ser Trp Pro Pro Phe Val Met Gly Ser Ala Phe Leu Val Ile  
195 200 205

Leu Leu Ile Met Lys Lys Leu Gly Lys Thr Asn Lys Lys Leu Arg Phe  
210 215 220

Leu Arg Ala Ser Gly Pro Leu Thr Ala Val Val Leu Gly Thr Leu Phe  
225 230 235 240

Val Lys Ile Phe Arg Pro Thr Ala Ile Ser Val Val Gly Glu Ile Pro  
245 250 255

Gln Gly Leu Pro Ser Phe Ser Ile Pro Arg Gly Phe Glu His Leu Met  
260 265 270

Ser Leu Met Pro Thr Ala Ile Leu Ile Thr Gly Val Ala Ile Leu Glu

275					280					285					
Ser	Val	Gly	Ile	Ala	Lys	Ala	Leu	Ala	Ala	Lys	Asn	Gly	Tyr	Glu	Leu
	290					295					300				
Asp	Ser	Asn	Lys	Glu	Leu	Phe	Gly	Leu	Gly	Leu	Ser	Asn	Ile	Cys	Gly
305					310					315					320
Ser	Phe	Phe	Ser	Ala	Tyr	Pro	Ala	Thr	Gly	Ser	Phe	Ser	Arg	Ser	Ala
				325					330					335	
Val	Asn	His	Glu	Ser	Gly	Ala	Lys	Thr	Gly	Leu	Ser	Gly	Ile	Ile	Met
			340					345					350		
Gly	Ile	Ile	Ile	Cys	Ser	Ala	Leu	Leu	Phe	Met	Thr	Pro	Leu	Phe	Thr
		355					360					365			
Asp	Ile	Pro	Gln	Cys	Ala	Leu	Ala	Ala	Ile	Val	Ile	Ser	Ala	Val	Thr
	370					375					380				
Gly	Leu	Val	Asp	Tyr	Glu	Glu	Ala	Ile	Phe	Leu	Trp	Gly	Ile	Asp	Lys
385					390					395					400
Lys	Asp	Phe	Phe	Leu	Trp	Ala	Met	Thr	Phe	Thr	Thr	Thr	Thr	Leu	Thr
				405					410					415	
Gly	Ile	Glu	Ile	Gly	Val	Leu	Val	Gly	Val	Gly	Phe	Ser	Leu	Ala	Phe
			420					425					430		
Val	Ile	His	Glu	Ser	Ala	Asn	Pro	His	Ile	Ala	Val	Leu	Gly	Arg	Leu
		435				440					445				
Pro	Gly	Thr	Thr	Val	Tyr	Arg	Asn	Thr	Leu	Gln	Tyr	Pro	Glu	Ala	Tyr
	450					455					460				
Thr	Tyr	Asn	Gly	Ile	Val	Val	Val	Arg	Val	Asp	Ala	Pro	Ile	Tyr	Phe
465					470					475					480
Ala	Asn	Ile	Ser	Tyr	Ile	Lys	Asp	Arg	Leu	Arg	Glu	Tyr	Glu	Leu	Lys
				485					490					495	
Leu	Pro	Asn	Ser	Asn	Arg	Gly	Pro	Asp	Val	Gly	Arg	Val	Tyr	Phe	Val
			500					505					510		
Ile	Leu	Glu	Met	Ser	Pro	Val	Thr	Tyr	Ile	Asp	Ser	Ser	Ala	Val	Gln
		515					520					525			
Ala	Leu	Lys	Asp	Leu	His	Gln	Glu	Tyr	Lys	Ala	Arg	Asp	Ile	Gln	Ile
	530					535					540				
Ala	Ile	Ala	Asn	Pro	Asn	Arg	Gln	Val	His	Leu	Leu	Leu	Ser	Arg	Ala
545					550					555					560
Gly	Ile	Ile	Asp	Met	Ile	Gly	Ala	Gly	Trp	Cys	Phe	Val	Arg	Val	His
				565					570					575	
Asp	Ala	Val	Gln	Val	Cys	Leu	Gln	His	Val	Arg	Ser	Ser	Ser	Ser	Asn
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Ala															

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 <211> 660  
 <212> PRT  
 <213> Sporobolus stapfianus

<400> 23

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			20					25					30		
Val	Glu	Val	His	Lys	Val	Val	Pro	Pro	Pro	Pro	Gln	Ser	Thr	Ala	Ser
		35					40					45			
Lys	Leu	Lys	Thr	Arg	Leu	Lys	Glu	Thr	Leu	Phe	Pro	Asp	Asp	Pro	Phe
	50					55					60				
Arg	Gly	Phe	Gln	Gly	Gln	Pro	Ala	Arg	Val	Gln	Trp	Val	Leu	Ala	Val
	65				70					75					80
Lys	Tyr	Leu	Phe	Pro	Ile	Leu	Asp	Trp	Leu	Pro	Ala	Tyr	Ser	Leu	Ser
				85					90					95	
Leu	Phe	Lys	Ser	Asp	Leu	Ile	Ala	Gly	Leu	Thr	Ile	Ala	Ser	Leu	Ala
			100					105					110		
Ile	Pro	Gln	Gly	Ile	Ser	Tyr	Ala	Lys	Leu	Ala	Asn	Leu	Pro	Pro	Leu
		115					120					125			
Ile	Gly	Leu	Tyr	Ser	Ser	Phe	Val	Pro	Pro	Leu	Val	Tyr	Ala	Val	Leu
	130					135					140				
Gly	Ser	Ser	Arg	Asp	Leu	Ala	Val	Gly	Pro	Val	Ser	Ile	Ser	Ser	Leu
145					150					155					160
Ile	Met	Gly	Pro	Cys	Cys	Ala	Ser	Arg	Gln	Pro	His	Cys	Gly	Ala	Asp
				165					170					175	
Ala	Val	Pro	Ala	Ala	Arg	Leu	His	Ala	Thr	Leu	Phe	Ala	Gly	Ile	Phe
			180					185					190		
Gln	Ala	Ser	Leu	Gly	Ile	Leu	Arg	Leu	Gly	Phe	Ile	Ile	Asp	Phe	Leu
		195					200					205			
Ser	Lys	Ala	Thr	Leu	Val	Gly	Phe	Met	Ala	Gly	Ala	Ala	Ile	Ile	Val
	210					215					220				
Ser	Leu	Gln	Gln	Leu	Lys	Ala	Leu	Leu	Gly	Ile	Val	His	Phe	Thr	Thr
225					230					235					240
Glu	Met	Gly	Ile	Val	Pro	Val	Met	Ala	Ser	Val	Phe	His	His	Thr	Lys
				245					250					255	
Glu	Trp	Ser	Trp	Gln	Thr	Ile	Leu	Met	Gly	Val	Cys	Phe	Leu	Val	Phe
			260					265					270		
Leu	Leu	Val	Ala	Arg	His	Val	Ser	Ile	Arg	Trp	Pro	Arg	Leu	Phe	Trp



275						280						285					
Val	Ser	Ala	Cys	Ala	Pro	Leu	Val	Ser	Val	Ile	Ile	Ser	Thr	Leu	Val		
290						295					300						
Val	Phe	Leu	Phe	Lys	Ala	Gln	Asn	His	Gly	Ile	Ser	Ile	Ile	Gly	Gln		
305					310					315					320		
Leu	Lys	Cys	Gly	Leu	Asn	Arg	Pro	Ser	Trp	Asp	Lys	Thr	Asn	Ile	Asp		
				325					330					335			
Thr	Thr	Tyr	Leu	Gly	Leu	Thr	Met	Lys	Thr	Gly	Leu	Val	Thr	Gly	Ile		
			340					345					350				
Ile	Ser	Leu	Thr	Glu	Gly	Ile	Ala	Val	Gly	Arg	Thr	Phe	Ala	Ser	Leu		
		355					360					365					
Lys	Glu	Tyr	Gln	Ile	Asp	Gly	Asn	Lys	Glu	Met	Met	Ala	Ile	Gly	Leu		
	370					375					380						
Met	Asn	Val	Val	Gly	Ser	Cys	Thr	Ser	Cys	Tyr	Val	Thr	Thr	Gly	Ala		
385					390					395					400		
Phe	Ser	Arg	Ser	Pro	Val	Asn	His	Asn	Ala	Gly	Cys	Lys	Thr	Ala	Met		
				405					410					415			
Ser	Asn	Val	Ile	Met	Ala	Leu	Thr	Val	Met	Val	Thr	Leu	Leu	Phe	Leu		
			420					425					430				
Met	Pro	Leu	Phe	Val	Tyr	Thr	Pro	Asn	Val	Val	Leu	Gly	Ala	Ile	Ile		
		435					440					445					
Ile	Ala	Ala	Val	Ile	Gly	Leu	Ile	Asp	Ile	Pro	Ala	Val	Tyr	His	Ile		
		450				455					460						
Trp	Lys	Met	Asp	Lys	Met	Asp	Phe	Leu	Val	Cys	Val	Cys	Ala	Phe	Ala		
465					470					475					480		
Gly	Val	Leu	Phe	Ile	Ser	Val	Gln	Glu	Gly	Leu	Ala	Ile	Ala	Val	Gly		
				485					490					495			
Ile	Ser	Val	Phe	Arg	Val	Leu	Leu	Gln	Ile	Thr	Arg	Pro	Lys	Ile	Thr		
			500					505					510				
Val	Gln	Gly	Asn	Ile	Met	Gly	Thr	Asp	Ile	Tyr	Arg	Asn	Leu	His	Gln		
		515					520					525					
Tyr	Lys	Asp	Ala	Gln	Arg	Ile	Pro	Gly	Phe	Leu	Ile	Leu	Ala	Thr	Glu		
	530					535					540						
Ala	Pro	Ile	Asn	Phe	Ala	Asn	Ser	Asn	Tyr	Leu	Asn	Glu	Arg	Ile	Lys		
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Arg	Trp	Ile	Glu	Glu	Glu	Ser	Ser	Ala	Gln	Thr	Lys	Gln	Thr	Glu	Leu		
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Arg	Phe	Val	Ile	Leu	Asp	Leu	Ser	Ala	Val	Pro	Ala	Ile	Asp	Thr	Ser		
			580					585					590				
Gly	Val	Ala	Phe	Leu	Ile	Asp	Ile	Lys	Lys	Ser	Ile	Glu	Lys	Arg	Gly		

595	600	605
Leu Glu Leu Val Leu Val Asn Pro Thr Gly Glu Gly His Gly Lys Asn		
610	615	620
Thr Ala Ser Glu Arg Gly Thr Gln Ala Phe Gln Val Gly Ile Ala Cys		
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Ile Leu Thr Thr Gly Glu Ala Val Ala Ser Leu Ser Ala Leu Ala Lys		
	645 650	655
Met Ala Ser Pro		
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 <213> Arabidopsis thaliana

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Ser Leu Gln Tyr Ser Val Lys Glu Thr Leu Phe Pro Asp Asp Pro Phe
35 40 45
Arg Gln Phe Lys Asn Gln Asn Ala Ser Arg Lys Phe Val Leu Gly Leu
50 55 60
Lys Tyr Phe Leu Pro Ile Phe Glu Trp Ala Pro Arg Tyr Asn Leu Lys
65 70 75 80
Phe Phe Lys Ser Asp Leu Ile Ala Gly Ile Thr Ile Ala Ser Leu Ala
85 90 95
Ile Pro Gln Gly Ile Ser Tyr Ala Lys Leu Ala Asn Leu Pro Pro Ile
100 105 110
Leu Gly Leu Tyr Ser Ser Phe Val Pro Pro Leu Val Tyr Ala Val Leu
115 120 125
Gly Ser Ser Arg Asp Leu Ala Val Gly Thr Val Ala Val Ala Ser Leu
130 135 140
Leu Thr Gly Ala Met Leu Ser Lys Glu Val Asp Ala Glu Lys Asp Pro
145 150 155 160
Lys Leu Tyr Leu His Leu Ala Phe Thr Ala Thr Phe Phe Ala Gly Val
165 170 175
Leu Glu Ala Ser Leu Gly Ile Phe Arg Leu Gly Phe Ile Val Asp Phe
180 185 190
Leu Ser His Ala Thr Ile Val Gly Phe Met Gly Gly Ala Ala Thr Val
195 200 205
Val Ser Leu Gln Gln Leu Lys Gly Ile Phe Gly Leu Lys His Phe Thr

210				215				220							
Asp	Ser	Thr	Asp	Val	Ile	Ser	Val	Met	Arg	Ser	Val	Phe	Ser	Gln	Thr
225					230					235					240
His	Glu	Trp	Arg	Trp	Glu	Ser	Gly	Val	Leu	Gly	Cys	Gly	Phe	Leu	Phe
				245					250					255	
Phe	Leu	Leu	Ser	Thr	Arg	Tyr	Phe	Ser	Ile	Lys	Lys	Pro	Lys	Phe	Phe
			260					265					270		
Trp	Val	Ala	Ala	Met	Ala	Pro	Leu	Thr	Ser	Val	Ile	Leu	Gly	Ser	Leu
		275					280						285		
Leu	Val	Tyr	Phe	Thr	His	Ala	Glu	Arg	His	Gly	Val	Gln	Val	Ile	Gly
		290				295					300				
Asp	Leu	Lys	Lys	Gly	Leu	Asn	Pro	Leu	Ser	Gly	Ser	Asp	Leu	Ile	Phe
305					310					315					320
Thr	Ser	Pro	Tyr	Met	Ser	Thr	Ala	Val	Lys	Thr	Gly	Leu	Ile	Thr	Gly
				325					330					335	
Ile	Ile	Ala	Leu	Ala	Glu	Gly	Ile	Ala	Val	Gly	Arg	Ser	Phe	Ala	Met
			340					345					350		
Phe	Lys	Asn	Tyr	Asn	Ile	Asp	Gly	Asn	Lys	Glu	Met	Ile	Ala	Phe	Gly
		355					360					365			
Met	Met	Asn	Ile	Val	Gly	Ser	Phe	Thr	Ser	Cys	Tyr	Leu	Thr	Thr	Gly
		370				375					380				
Pro	Phe	Ser	Arg	Ser	Ala	Val	Asn	Tyr	Asn	Ala	Gly	Cys	Lys	Thr	Ala
385					390					395					400
Met	Ser	Asn	Ile	Val	Met	Ala	Ile	Ala	Val	Met	Phe	Thr	Leu	Leu	Phe
			405						410					415	
Leu	Thr	Pro	Leu	Phe	His	Tyr	Thr	Pro	Leu	Val	Val	Leu	Ser	Ala	Ile
			420					425					430		
Ile	Ile	Ser	Ala	Met	Leu	Gly	Leu	Ile	Asp	Tyr	Gln	Ala	Ala	Ile	His
		435					440					445			
Leu	Trp	Lys	Val	Asp	Lys	Phe	Asp	Phe	Leu	Val	Cys	Met	Ser	Ala	Tyr
		450				455					460				
Val	Gly	Val	Val	Phe	Gly	Ser	Val	Glu	Ile	Gly	Leu	Val	Val	Ala	Val
465					470					475					480
Ala	Ile	Ser	Ile	Ala	Arg	Leu	Leu	Leu	Phe	Val	Ser	Arg	Pro	Lys	Thr
			485						490					495	
Ala	Val	Lys	Gly	Asn	Ile	Pro	Asn	Ser	Met	Ile	Tyr	Arg	Asn	Thr	Glu
			500					505					510		
Gln	Tyr	Pro	Ser	Ser	Arg	Thr	Val	Pro	Gly	Ile	Leu	Ile	Leu	Glu	Ile
		515					520						525		
Asp	Ala	Pro	Ile	Tyr	Phe	Ala	Asn	Ala	Ser	Tyr	Leu	Arg	Glu	Arg	Ile

530					535					540					
Ile	Arg	Trp	Ile	Asp	Glu	Glu	Glu	Glu	Arg	Val	Lys	Gln	Ser	Gly	Glu
545					550					555					560
Ser	Ser	Leu	Gln	Tyr	Ile	Ile	Leu	Asp	Met	Ser	Ala	Val	Gly	Asn	Ile
				565					570					575	
Asp	Thr	Ser	Gly	Ile	Ser	Met	Met	Val	Glu	Ile	Lys	Lys	Val	Ile	Asp
			580					585					590		
Arg	Arg	Ala	Leu	Lys	Leu	Val	Leu	Ser	Asn	Pro	Lys	Gly	Glu	Val	Val
		595					600					605			
Lys	Lys	Leu	Thr	Arg	Ser	Lys	Phe	Ile	Gly	Asp	His	Leu	Gly	Lys	Glu
	610					615					620				
Trp	Met	Phe	Leu	Thr	Val	Gly	Glu	Ala	Val	Glu	Ala	Cys	Ser	Tyr	Met
625					630					635					640
Leu	His	Thr	Phe	Lys	Thr	Glu	Pro	Ala	Ser	Lys	Asn	Glu	Pro	Trp	Asn
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Asn Val

<210> 25  
 <211> 644  
 <212> PRT  
 <213> Stylosanthes hamata

<400> 25															
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Leu	Asn	Ser	Pro	Asn	Pro	Pro	Pro	Leu	Thr	Lys	Lys	Phe	Leu	Gly	Pro
			20					25					30		
Leu	Lys	Asp	Asn	Lys	Phe	Phe	Thr	Ser	Ser	Ser	Ser	Lys	Lys	Glu	Thr
		35					40					45			
Arg	Ala	Val	Ser	Phe	Leu	Ala	Ser	Leu	Phe	Pro	Ile	Leu	Ser	Trp	Ile
	50					55					60				
Arg	Thr	Tyr	Ser	Ala	Thr	Lys	Phe	Lys	Asp	Asp	Leu	Leu	Ser	Gly	Leu
	65				70					75					80
Thr	Leu	Ala	Ser	Leu	Ser	Ile	Pro	Gln	Ser	Ile	Gly	Tyr	Ala	Asn	Leu
				85					90					95	
Ala	Lys	Leu	Asp	Pro	Gln	Tyr	Gly	Leu	Tyr	Thr	Ser	Val	Ile	Pro	Pro
			100					105					110		
Val	Ile	Tyr	Ala	Leu	Met	Gly	Ser	Ser	Arg	Glu	Ile	Ala	Ile	Gly	Pro
		115					120					125			
Val	Ala	Val	Val	Ser	Met	Leu	Leu	Ser	Ser	Leu	Val	Pro	Lys	Val	Ile
	130					135					140				
Asp	Pro	Asp	Ala	His	Pro	Asn	Asp	Tyr	Arg	Asn	Leu	Val	Phe	Thr	Val

145					150					155					160
Thr	Leu	Phe	Ala	Gly	Ile	Phe	Gln	Thr	Ala	Phe	Gly	Val	Leu	Arg	Leu
				165					170					175	
Gly	Phe	Leu	Val	Asp	Phe	Leu	Ser	His	Ala	Ala	Leu	Val	Gly	Phe	Met
			180					185					190		
Ala	Gly	Ala	Ala	Ile	Val	Ile	Gly	Leu	Gln	Gln	Leu	Lys	Gly	Leu	Leu
			195				200					205			
Gly	Leu	Thr	His	Phe	Thr	Thr	Lys	Thr	Asp	Ala	Val	Ala	Val	Leu	Lys
	210					215					220				
Ser	Val	Tyr	Thr	Ser	Leu	His	Gln	Gln	Ile	Thr	Ser	Ser	Glu	Asn	Trp
225					230					235					240
Ser	Pro	Leu	Asn	Phe	Val	Ile	Gly	Cys	Ser	Phe	Leu	Ile	Phe	Leu	Leu
				245					250					255	
Ala	Ala	Arg	Phe	Ile	Gly	Arg	Arg	Asn	Lys	Lys	Phe	Phe	Trp	Leu	Pro
			260					265					270		
Ala	Ile	Ala	Pro	Leu	Leu	Ser	Val	Ile	Leu	Ser	Thr	Leu	Ile	Val	Phe
		275					280					285			
Leu	Ser	Lys	Gly	Asp	Lys	His	Gly	Val	Asn	Ile	Ile	Lys	His	Val	Gln
	290					295					300				
Gly	Gly	Leu	Asn	Pro	Ser	Ser	Val	His	Lys	Leu	Gln	Leu	Asn	Gly	Pro
305					310					315					320
His	Val	Gly	Gln	Ala	Ala	Lys	Ile	Gly	Leu	Ile	Ser	Ala	Ile	Ile	Ala
				325					330					335	
Leu	Thr	Glu	Ala	Ile	Ala	Val	Gly	Arg	Ser	Phe	Ala	Asn	Ile	Lys	Gly
			340					345					350		
Tyr	His	Leu	Asp	Gly	Asn	Lys	Glu	Met	Leu	Ala	Met	Gly	Cys	Met	Asn
		355					360					365			
Ile	Ala	Gly	Ser	Leu	Thr	Ser	Cys	Tyr	Val	Ser	Thr	Gly	Ser	Phe	Ser
	370					375					380				
Arg	Thr	Ala	Val	Asn	Phe	Ser	Ala	Gly	Cys	Lys	Thr	Ala	Val	Ser	Asn
385					390					395					400
Ile	Val	Met	Ala	Val	Thr	Val	Leu	Leu	Cys	Leu	Glu	Leu	Phe	Thr	Arg
				405					410					415	
Leu	Leu	Tyr	Tyr	Thr	Pro	Met	Ala	Ile	Leu	Ala	Ser	Ile	Ile	Leu	Ser
			420					425					430		
Ala	Leu	Pro	Gly	Leu	Ile	Asp	Ile	Gly	Glu	Ala	Tyr	His	Ile	Trp	Lys
		435					440					445			
Val	Asp	Lys	Phe	Asp	Phe	Leu	Ala	Cys	Leu	Gly	Ala	Phe	Phe	Gly	Val
	450					455					460				
Leu	Phe	Val	Ser	Ile	Glu	Ile	Gly	Leu	Leu	Ile	Ala	Leu	Ser	Ile	Ser

465		470		475		480									
Phe	Ala	Lys	Ile	Leu	Leu	Gln	Ala	Ile	Arg	Pro	Gly	Val	Glu	Val	Leu
				485					490					495	
Gly	Arg	Ile	Pro	Thr	Thr	Glu	Ala	Tyr	Cys	Asp	Val	Ala	Gln	Tyr	Pro
			500					505					510		
Met	Ala	Val	Thr	Thr	Pro	Gly	Ile	Leu	Val	Ile	Arg	Ile	Ser	Ser	Gly
		515					520					525			
Ser	Leu	Cys	Phe	Ala	Asn	Ala	Gly	Phe	Val	Arg	Glu	Arg	Ile	Leu	Lys
	530					535					540				
Trp	Val	Glu	Asp	Glu	Glu	Gln	Asp	Asn	Ile	Glu	Glu	Ala	Ala	Lys	Gly
545					550					555					560
Arg	Val	Gln	Ala	Ile	Ile	Ile	Asp	Met	Thr	Asp	Leu	Thr	Asn	Val	Asp
				565					570					575	
Thr	Ser	Gly	Ile	Leu	Ala	Leu	Glu	Glu	Leu	His	Lys	Lys	Leu	Leu	Ser
			580					585					590		
Arg	Gly	Val	Glu	Leu	Ala	Met	Val	Asn	Pro	Arg	Trp	Glu	Val	Ile	His
		595					600					605			
Lys	Leu	Lys	Val	Ala	Asn	Phe	Val	Asp	Lys	Ile	Gly	Lys	Glu	Arg	Val
	610					615					620				
Phe	Leu	Thr	Val	Ala	Glu	Ala	Val	Asp	Ala	Cys	Leu	Ser	Ser	Arg	Phe
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Ala Asn Ser Ala

<210> 26  
 <211> 646  
 <212> PRT  
 <213> Arabidopsis thaliana

<400> 26

Met	Gly	Thr	Glu	Asp	Tyr	Thr	Phe	Pro	Gln	Gly	Ala	Glu	Glu	Leu	His
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Arg	Arg	His	His	Thr	Val	Glu	Ala	Pro	Gln	Pro	Gln	Pro	Phe	Leu	Lys
			20					25					30		
Ser	Leu	Gln	Tyr	Ser	Val	Lys	Glu	Thr	Leu	Phe	Pro	Asp	Asp	Pro	Phe
		35					40					45			
Arg	Gln	Phe	Lys	Asn	Gln	Asn	Ala	Ser	Arg	Lys	Phe	Val	Leu	Gly	Leu
	50					55					60				
Lys	Tyr	Phe	Leu	Pro	Ile	Phe	Glu	Trp	Ala	Pro	Arg	Tyr	Asn	Leu	Lys
65					70					75				80	
Phe	Phe	Lys	Ser	Asp	Leu	Ile	Ala	Gly	Ile	Thr	Ile	Ala	Ser	Leu	Ala
				85					90					95	

Ile Pro Gln Gly Ile Ser Tyr Ala Lys Leu Ala Asn Leu Pro Pro Ile

100						105						110					
Leu	Gly	Leu	Tyr	Ser	Ser	Phe	Val	Pro	Pro	Leu	Val	Tyr	Ala	Val	Leu		
		115					120					125					
Gly	Ser	Ser	Arg	Asp	Leu	Ala	Val	Gly	Thr	Val	Ala	Val	Ala	Ser	Leu		
	130					135					140						
Leu	Thr	Gly	Ala	Met	Leu	Ser	Lys	Glu	Val	Asp	Ala	Glu	Lys	Asp	Pro		
145					150					155					160		
Lys	Leu	Tyr	Leu	His	Leu	Ala	Phe	Thr	Ala	Thr	Phe	Phe	Ala	Gly	Val		
				165					170					175			
Leu	Glu	Ala	Ser	Leu	Gly	Ile	Phe	Arg	Leu	Gly	Phe	Ile	Val	Asp	Phe		
			180					185					190				
Leu	Ser	His	Ala	Thr	Ile	Val	Gly	Phe	Met	Gly	Gly	Ala	Ala	Thr	Val		
	195						200					205					
Val	Ser	Leu	Gln	Gln	Leu	Lys	Gly	Ile	Phe	Gly	Leu	Lys	His	Phe	Thr		
	210					215					220						
Asp	Ser	Thr	Asp	Val	Ile	Ser	Val	Met	Arg	Ser	Val	Phe	Ser	Gln	Thr		
225					230					235					240		
His	Glu	Trp	Arg	Trp	Glu	Ser	Gly	Val	Leu	Gly	Cys	Gly	Phe	Leu	Phe		
				245					250					255			
Phe	Leu	Leu	Ser	Thr	Arg	Tyr	Phe	Ser	Ile	Lys	Lys	Pro	Lys	Phe	Phe		
			260					265					270				
Trp	Val	Ala	Ala	Met	Ala	Pro	Leu	Thr	Ser	Val	Ile	Leu	Gly	Ser	Leu		
		275					280					285					
Leu	Val	Tyr	Phe	Thr	His	Ala	Glu	Arg	His	Gly	Val	Gln	Val	Gly	Ser		
	290					295					300						
Asp	Leu	Ile	Phe	Thr	Ser	Pro	Tyr	Met	Ser	Thr	Ala	Val	Lys	Thr	Gly		
305					310					315					320		
Leu	Ile	Thr	Gly	Ile	Ile	Ala	Leu	Ala	Glu	Gly	Val	Ala	Val	Gly	Arg		
				325					330					335			
Ser	Phe	Ala	Met	Phe	Lys	Asn	Tyr	Asn	Ile	Asp	Gly	Asn	Lys	Glu	Met		
			340					345					350				
Ile	Ala	Phe	Gly	Met	Met	Asn	Ile	Val	Gly	Ser	Phe	Thr	Ser	Cys	Tyr		
		355					360					365					
Leu	Thr	Thr	Gly	Pro	Phe	Ser	Arg	Ser	Ala	Val	Asn	Tyr	Asn	Ala	Gly		
	370					375					380						
Cys	Lys	Thr	Ala	Met	Ser	Asn	Ile	Val	Met	Ala	Ile	Ala	Val	Met	Phe		
385					390					395					400		
Thr	Leu	Leu	Phe	Leu	Thr	Pro	Leu	Phe	His	Tyr	Thr	Pro	Leu	Val	Val		
				405					410					415			
Leu	Ser	Ala	Ile	Ile	Ile	Ser	Ala	Met	Leu	Gly	Leu	Ile	Asp	Tyr	Gln		

420					425					430					
Ala	Ala	Ile	His	Leu	Trp	Lys	Val	Asp	Lys	Phe	Asp	Phe	Leu	Val	Cys
		435					440					445			
Met	Ser	Ala	Tyr	Val	Gly	Val	Val	Phe	Gly	Ser	Val	Glu	Ile	Gly	Leu
	450					455					460				
Val	Val	Ala	Val	Ala	Ile	Ser	Ile	Ala	Arg	Leu	Leu	Leu	Phe	Val	Ser
	465					470					475				480
Arg	Pro	Lys	Thr	Ala	Val	Lys	Gly	Asn	Ile	Pro	Asn	Ser	Met	Ile	Tyr
				485					490					495	
Arg	Asn	Thr	Glu	Gln	Tyr	Pro	Ser	Ser	Arg	Thr	Val	Pro	Gly	Ile	Leu
			500					505					510		
Ile	Leu	Glu	Ile	Asp	Ala	Pro	Ile	Tyr	Phe	Ala	Asn	Ala	Ser	Tyr	Leu
		515					520					525			
Arg	Glu	Arg	Ile	Ile	Arg	Trp	Ile	Asp	Glu	Glu	Glu	Glu	Arg	Val	Lys
	530					535						540			
Gln	Ser	Gly	Glu	Ser	Ser	Leu	Gln	Tyr	Ile	Ile	Leu	Asp	Met	Ser	Ala
	545					550					555				560
Val	Gly	Asn	Ile	Asp	Thr	Ser	Gly	Ile	Ser	Met	Met	Val	Glu	Ile	Lys
				565					570					575	
Lys	Val	Ile	Asp	Arg	Arg	Ala	Leu	Lys	Leu	Val	Leu	Ser	Asn	Pro	Lys
			580					585					590		
Gly	Glu	Val	Val	Lys	Lys	Leu	Thr	Arg	Ser	Lys	Phe	Ile	Gly	Asp	His
		595					600					605			
Leu	Gly	Lys	Glu	Trp	Met	Phe	Leu	Thr	Val	Gly	Glu	Ala	Val	Glu	Ala
	610					615					620				
Cys	Ser	Tyr	Met	Leu	His	Thr	Phe	Lys	Thr	Glu	Pro	Ala	Ser	Lys	Asn
	625					630					635				640
Glu	Pro	Trp	Asn	Asn	Val										
				645											

<210> 27  
 <211> 233  
 <212> PRT  
 <213> Zea mays

<400> 27  
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 Gln Asn Glu Ile Asp Pro Lys Thr His Pro Leu Glu Tyr Arg Arg Leu  
 20 25 30  
 Ala Phe Thr Ala Thr Phe Phe Ala Gly Val Thr Gln Ala Ala Leu Gly  
 35 40 45  
 Phe Phe Arg Leu Gly Phe Ile Ile Glu Phe Leu Ser His Ala Ala Ile



50					55					60					
Val	Gly	Phe	Met	Ala	Gly	Ala	Ala	Ile	Thr	Ile	Ala	Leu	Gln	Gln	Leu
65					70					75					80
Lys	Gly	Phe	Leu	Gly	Ile	Ala	Asn	Phe	Thr	Lys	Lys	Ser	Asp	Ile	Val
				85					90					95	
Ser	Val	Met	Lys	Ser	Val	Trp	Gly	Asn	Val	His	His	Gly	Trp	Asn	Trp
			100					105					110		
Gln	Thr	Ile	Leu	Ile	Gly	Ala	Thr	Phe	Leu	Ala	Phe	Leu	Leu	Val	Ala
		115					120					125			
Lys	Tyr	Ile	Gly	Lys	Arg	Asn	Lys	Lys	Leu	Phe	Trp	Val	Ser	Ala	Ile
	130					135					140				
Ala	Pro	Leu	Thr	Ser	Val	Ile	Ile	Ser	Thr	Phe	Phe	Val	Tyr	Ile	Thr
145					150					155					160
Arg	Ala	Asp	Lys	His	Gly	Val	Ala	Ile	Val	Lys	Asn	Ile	Arg	Lys	Gly
				165					170					175	
Ile	Asn	Pro	Pro	Ser	Ala	Ser	Leu	Ile	Tyr	Phe	Thr	Gly	Pro	Tyr	Leu
			180					185					190		
Ala	Thr	Gly	Phe	Lys	Ile	Gly	Ile	Val	Ala	Gly	Met	Ile	Gly	Leu	Thr
		195					200					205			
Glu	Ala	Ile	Ala	Ile	Gly	Arg	Thr	Phe	Ala	Ala	Leu	Lys	Asp	Tyr	Arg
	210					215					220				
Ile	Asp	Gly	Asn	Lys	Glu	Met	Val	Ala							
225					230										

<210> 28  
 <211> 646  
 <212> PRT  
 <213> Arabidopsis thaliana

<400> 28															
Met	Ser	Ser	Lys	Arg	Ala	Ser	Gln	Tyr	His	Gln	Val	Glu	Ile	Pro	Pro
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Pro	Gln	Pro	Phe	Leu	Lys	Ser	Leu	Lys	Asn	Thr	Leu	Asn	Glu	Ile	Leu
			20					25					30		
Phe	Ala	Asp	Asp	Pro	Phe	Arg	Arg	Ile	Arg	Asn	Glu	Ser	Lys	Thr	Ser
		35					40					45			
Lys	Lys	Ile	Glu	Leu	Gly	Leu	Arg	His	Val	Phe	Pro	Ile	Leu	Glu	Trp
	50					55					60				
Ala	Arg	Gly	Tyr	Ser	Leu	Glu	Tyr	Leu	Lys	Ser	Asp	Val	Ile	Ser	Gly
65					70					75					80
Ile	Thr	Ile	Ala	Ser	Leu	Ala	Ile	Pro	Gln	Gly	Ile	Ser	Tyr	Ala	Gln
				85					90					95	
Leu	Ala	Asn	Leu	Pro	Pro	Ile	Leu	Gly	Leu	Tyr	Ser	Ser	Leu	Val	Pro

100					105					110					
Pro	Leu	Val	Tyr	Ala	Ile	Met	Gly	Ser	Ser	Arg	Asp	Leu	Ala	Val	Gly
		115					120					125			
Thr	Val	Ala	Val	Ala	Ser	Leu	Leu	Thr	Ala	Ala	Met	Leu	Gly	Lys	Glu
	130					135					140				
Val	Asn	Ala	Val	Val	Asn	Pro	Lys	Leu	Tyr	Leu	His	Leu	Ala	Phe	Thr
	145					150					155				160
Ala	Thr	Phe	Phe	Ala	Gly	Leu	Met	Gln	Thr	Cys	Leu	Gly	Leu	Leu	Arg
				165					170						175
Leu	Gly	Phe	Val	Val	Glu	Ile	Leu	Ser	His	Ala	Ala	Ile	Val	Gly	Phe
			180					185					190		
Met	Gly	Gly	Ala	Ala	Thr	Val	Val	Cys	Leu	Gln	Gln	Leu	Lys	Gly	Leu
		195					200					205			
Leu	Gly	Leu	His	His	Phe	Thr	His	Ser	Thr	Asp	Ile	Val	Thr	Val	Leu
	210					215					220				
Arg	Ser	Ile	Phe	Ser	Gln	Ser	His	Met	Trp	Arg	Trp	Glu	Ser	Gly	Val
	225					230					235				240
Leu	Gly	Cys	Cys	Phe	Leu	Ile	Phe	Leu	Leu	Thr	Thr	Lys	Tyr	Ile	Ser
				245					250					255	
Lys	Lys	Arg	Pro	Lys	Leu	Phe	Trp	Ile	Ser	Ala	Met	Ser	Pro	Leu	Val
			260					265					270		
Ser	Val	Ile	Phe	Gly	Thr	Ile	Phe	Leu	Tyr	Phe	Leu	His	Asp	Gln	Phe
		275					280					285			
His	Gly	Ile	Gln	Phe	Ile	Gly	Glu	Leu	Lys	Lys	Gly	Ile	Asn	Pro	Pro
	290					295					300				
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Lys	Val	Gly	Ile	Ile	Thr	Gly	Val	Ile	Ala	Leu	Ala	Glu	Gly	Ile	Ala
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Val	Gly	Arg	Ser	Phe	Ala	Met	Tyr	Lys	Asn	Tyr	Asn	Ile	Asp	Gly	Asn
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Lys	Glu	Met	Ile	Ala	Phe	Gly	Met	Met	Asn	Ile	Leu	Gly	Ser	Phe	Ser
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Ser	Cys	Tyr	Leu	Thr	Thr	Gly	Pro	Phe	Ser	Arg	Ser	Ala	Val	Asn	Tyr
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Asn	Ala	Gly	Cys	Lys	Thr	Ala	Leu	Ser	Asn	Val	Val	Met	Ala	Val	Ala
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Val	Ala	Val	Thr	Leu	Leu	Phe	Leu	Thr	Pro	Leu	Phe	Phe	Tyr	Thr	Pro
				405					410					415	
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Asp	Tyr	Glu	Ala	Ala	Ile	His	Leu	Trp	Lys	Leu	Asp	Lys	Phe	Asp	Phe
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Ile	Gly	Leu	Ile	Leu	Ser	Val	Gly	Ile	Ser	Val	Met	Arg	Leu	Val	Leu
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Phe	Val	Gly	Arg	Pro	Lys	Ile	Tyr	Val	Met	Gly	Asn	Ile	Gln	Asn	Ser
				485					490					495	
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			500					505					510		
Ser	Leu	Leu	Ile	Leu	His	Ile	Asp	Gly	Pro	Ile	Tyr	Phe	Ala	Asn	Ser
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Thr	Tyr	Leu	Arg	Asp	Arg	Ile	Gly	Arg	Trp	Ile	Asp	Glu	Glu	Glu	Asp
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Lys	Leu	Arg	Thr	Ser	Gly	Asp	Ile	Ser	Leu	Gln	Tyr	Ile	Val	Leu	Asp
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Met	Ser	Ala	Val	Gly	Asn	Ile	Asp	Thr	Ser	Gly	Ile	Ser	Met	Leu	Glu
				565					570					575	
Glu	Leu	Asn	Lys	Ile	Leu	Gly	Arg	Arg	Glu	Leu	Lys	Leu	Val	Ile	Ala
		580						585					590		
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Ala	Ala	Cys	Asp	Phe	Met	Leu	His	Thr	Ala	Lys	Pro	Asp	Ser	Pro	Val
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 Gln Tyr Ile Phe Pro Ile Leu Gln Trp Cys Pro Glu Tyr Ser Phe Ser

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Ile	Pro	Gln	Gly	Ile 85	Ser	Tyr	Ala	Asn	Val 90	Ala	Asn	Leu	Pro	Pro	Ile 95
Val	Gly	Leu	Tyr 100	Ser	Ser	Phe	Val	Pro 105	Pro	Leu	Val	Tyr	Ala	Val	Leu 110
Gly	Ser	Ser	Arg	Asp	Leu	Ala	Val 120	Gly	Pro	Val	Ser	Ile	Ala	Ser	Leu 125
Ile	Leu 130	Gly	Ser	Met	Leu	Arg 135	Gln	Gln	Val	Ser	Pro	Val	Asp	Asp	Pro 140
Val 145	Leu	Phe	Leu	Gln	Leu	Ala	Phe	Ser	Ser	Thr 155	Phe	Phe	Ala	Gly	Leu 160
Phe	Gln	Ala	Ser	Leu 165	Gly	Ile	Leu	Arg	Leu	Gly 170	Phe	Ile	Ile	Asp	Phe 175
Leu	Ser	Lys	Ala 180	Thr	Leu	Ile	Gly	Phe	Met	Gly	Gly	Ala	Ala	Ile	Ile 190
Val	Ser	Leu 195	Gln	Gln	Leu	Lys	Gly 200	Leu	Leu	Gly	Ile	Thr	His	Phe	Thr 205
Lys	His 210	Met	Ser	Val	Val	Pro 215	Val	Leu	Ser	Ser	Val	Phe	Gln	His	Thr 220
Asn 225	Glu	Trp	Ser	Trp	Gln	Thr	Ile	Val	Met	Gly 235	Val	Cys	Phe	Leu	Leu 240
Phe	Leu	Leu	Ser	Thr 245	Arg	His	Leu	Ser	Met	Lys	Lys	Pro	Lys	Leu	Phe 255
Trp	Val	Ser	Ala 260	Gly	Ala	Pro	Leu	Leu	Ser	Val	Ile	Val	Ser	Thr	Leu 270
Leu	Val	Phe 275	Val	Phe	Arg	Ala	Glu 280	Arg	His	Gly	Ile	Ser	Val	Ile	Gly 285
Lys	Leu 290	Pro	Glu	Gly	Leu	Asn 295	Pro	Pro	Ser	Trp	Asn	Met	Leu	Gln	Phe 300
His 305	Gly	Ser	His	Leu	Ala	Leu	Val	Ala	Lys	Thr 315	Gly	Leu	Val	Thr	Gly 320
Ile	Val	Ser	Leu	Thr 325	Glu	Gly	Ile	Ala	Val 330	Gly	Arg	Thr	Phe	Ala	Ala 335
Leu	Lys	Asn	Tyr 340	His	Val	Asp	Gly	Asn 345	Lys	Glu	Met	Ile	Ala	Ile	Gly 350
Leu	Met	Asn	Val	Val	Gly	Ser	Ala 360	Thr	Ser	Cys	Tyr	Val	Thr	Thr	Gly 365
Ala	Phe	Ser	Arg	Ser	Ala	Val	Asn	Asn	Asn	Ala	Gly	Ala	Lys	Thr	Ala

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Leu	Met	Pro	Leu	Phe	Glu	Tyr	Thr	Pro	Asn	Val	Val	Leu	Gly	Ala	Ile
				405					410					415	
Ile	Val	Thr	Ala	Val	Ile	Gly	Leu	Ile	Asp	Leu	Pro	Ala	Ala	Cys	His
			420					425					430		
Ile	Trp	Lys	Ile	Asp	Lys	Phe	Asp	Phe	Leu	Val	Met	Leu	Cys	Ala	Phe
		435					440					445			
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	450					455					460				
Gly	Leu	Ser	Leu	Phe	Lys	Ile	Leu	Met	Gln	Val	Thr	Arg	Pro	Lys	Met
465				470						475					480
Val	Ile	Met	Gly	Asn	Ile	Pro	Gly	Thr	Asp	Ile	Tyr	Arg	Asp	Leu	His
				485					490					495	
His	Tyr	Lys	Glu	Ala	Gln	Arg	Ile	Pro	Gly	Phe	Leu	Val	Leu	Ser	Ile
			500					505					510		
Glu	Ser	Pro	Val	Asn	Phe	Ala	Asn	Ser	Asn	Tyr	Leu	Thr	Glu	Arg	Thr
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Ser	Arg	Trp	Ile	Glu	Glu	Cys	Glu	Glu	Glu	Glu	Ala	Gln	Glu	Lys	His
	530					535					540				
Ser	Ser	Leu	Gln	Phe	Leu	Ile	Leu	Glu	Met	Ser	Ala	Val	Ser	Gly	Val
545				550					555						560
Asp	Thr	Asn	Gly	Val	Ser	Phe	Phe	Lys	Glu	Leu	Lys	Lys	Thr	Thr	Ala
			565					570						575	
Lys	Lys	Asp	Ile	Glu	Leu	Val	Phe	Val	Asn	Pro	Leu	Ser	Glu	Val	Val
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Cys Ser Gln Thr Ala Ser Gln Arg His Thr Asp Ser Thr His His His															

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Phe	Ala	Glu	Gly	Val	Lys	Glu	Thr	Phe	Phe	Ala	Asp	Asp	Pro	Leu	Arg
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Glu	Tyr	Lys	Asp	Gln	Pro	Arg	Ser	Lys	Lys	Leu	Trp	Leu	Ser	Leu	Val
	65					70					75				80
His	Leu	Phe	Pro	Val	Leu	Asp	Trp	Ser	Arg	Ser	Tyr	Thr	Phe	Gly	Lys
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Phe	Lys	Gly	Asp	Leu	Val	Ala	Gly	Leu	Thr	Ile	Ala	Ser	Leu	Cys	Ile
			100					105					110		
Pro	Gln	Asp	Ile	Gly	Tyr	Ala	Lys	Leu	Ala	Asn	Leu	Gln	Pro	His	Val
		115					120					125			
Gly	Leu	Tyr	Ser	Ser	Phe	Val	Pro	Pro	Leu	Ile	Tyr	Ala	Leu	Met	Gly
	130					135					140				
Ser	Ser	Arg	Asp	Ile	Ala	Ile	Gly	Pro	Val	Ala	Val	Val	Ser	Leu	Leu
	145					150					155				160
Leu	Gly	Thr	Leu	Leu	Gln	Glu	Glu	Ile	Asp	Pro	Val	Lys	Asn	Pro	Leu
				165					170					175	
Glu	Tyr	Ser	Arg	Leu	Ala	Phe	Thr	Ala	Thr	Phe	Phe	Ala	Gly	Ile	Thr
			180					185					190		
Gln	Ala	Met	Leu	Gly	Phe	Phe	Arg	Leu	Gly	Phe	Ile	Ile	Glu	Phe	Leu
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Ala	Leu	Gln	Gln	Leu	Lys	Gly	Leu	Leu	Gly	Ile	Ala	Lys	Phe	Thr	Lys
	225					230					235				240
Lys	Ser	Asp	Ile	Ile	Ser	Val	Met	Glu	Ser	Val	Trp	Gly	Asn	Val	Gln
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His	Gly	Trp	Asn	Trp	Gln	Thr	Ile	Leu	Ile	Gly	Ser	Ser	Phe	Leu	Ala
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Phe	Leu	Leu	Thr	Thr	Lys	Tyr	Ile	Ala	Lys	Lys	Asn	Lys	Lys	Leu	Phe
			275				280					285			
Trp	Val	Ser	Ala	Ile	Ala	Pro	Leu	Ile	Ser	Val	Val	Ile	Ser	Thr	Phe
	290					295					300				
Cys	Val	Tyr	Ile	Thr	Arg	Ala	Asp	Lys	Gln	Gly	Val	Ala	Ile	Val	Lys
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Asn	Ile	Lys	Gln	Gly	Ile	Asn	Pro	Pro	Ser	Phe	Asp	Leu	Ile	Tyr	Trp
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Met	Val	Ala	Leu	Thr	Glu	Ala	Ile	Ala	Ile	Gly	Arg	Thr	Phe	Ala	Ala
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Thr	Met	Asn	Ile	Val	Gly	Ser	Met	Thr	Ser	Cys	Tyr	Val	Ala	Thr	Gly
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Ser	Phe	Ser	Arg	Ser	Ala	Val	Asn	Tyr	Met	Ala	Gly	Cys	Lys	Thr	Ala
				405					410					415	
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			420					425					430		
Ile	Thr	Pro	Leu	Phe	Lys	Tyr	Thr	Pro	Asn	Ala	Ile	Leu	Ala	Ser	Ile
		435					440					445			
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	450					455					460				
Ile	Trp	Lys	Val	Asp	Lys	Met	Asp	Phe	Val	Ala	Leu	Leu	Gly	Ala	Phe
465					470					475					480
Phe	Gly	Val	Val	Phe	Ala	Ser	Val	Glu	Tyr	Gly	Leu	Leu	Ile	Ala	Val
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			500					505					510		
Ala	Leu	Leu	Gly	Asn	Leu	Pro	Arg	Thr	Thr	Ile	Tyr	Arg	Asn	Val	Glu
			515				520					525			
Gln	Tyr	Pro	Glu	Ala	Thr	Lys	Val	Pro	Gly	Val	Met	Ile	Val	Arg	Val
	530					535					540				
Asp	Ser	Ala	Ile	Tyr	Phe	Thr	Asn	Ser	Asn	Tyr	Val	Lys	Glu	Arg	Ile
545					550					555					560
Leu	Arg	Trp	Leu	Arg	Asp	Glu	Glu	Glu	Gln	Gln	Gln	Glu	Gln	Lys	Leu
				565					570					575	
Ser	Lys	Thr	Glu	Phe	Leu	Ile	Val	Glu	Leu	Ser	Pro	Val	Thr	Asp	Ile
			580					585					590		
Asp	Thr	Ser	Gly	Ile	His	Ala	Leu	Glu	Glu	Leu	Leu	Lys	Ala	Leu	Glu
		595					600					605			
Lys	Arg	Lys	Ile	Gln	Leu	Ile	Leu	Ala	Asn	Pro	Gly	Pro	Ala	Val	Ile
	610					615					620				
Gln	Lys	Leu	Arg	Ser	Ala	Lys	Phe	Thr	Asp	Leu	Ile	Gly	Asp	Asp	Lys
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 Ala Lys Ile Lys Arg Met Arg Leu Val Asp Trp Ile Asp Thr Leu Phe  
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 Tyr Ser Ser Phe Val Pro Val Phe Val Tyr Ala Ile Phe Gly Ser Ser  
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 Asn Ala Leu Gly Gly Ile Ala Asp Thr Asn Glu Glu Leu His Ile Glu  
                   165                  170                  175  
 Leu Ala Ile Leu Leu Ala Leu Leu Val Gly Ile Leu Glu Cys Ile Met  
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 Gly Leu Leu Arg Leu Gly Trp Leu Ile Arg Phe Ile Ser His Ser Val  
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 Ile Lys Tyr Phe Leu Gly Tyr Ser Ile Ala Arg Ser Ser Lys Ile Val  
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 Pro Ile Val Glu Ser Ile Ile Ala Gly Ala Asp Lys Phe Gln Trp Pro  
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 Pro Phe Val Met Gly Ser Leu Ile Leu Val Ile Leu Gln Val Met Lys  
                   260                  265                  270  
 His Val Gly Lys Ala Lys Lys Glu Leu Gln Phe Leu Arg Ala Ala Ala



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Phe	Ser	Phe	Pro	Arg	Ser	Phe	Asp	His	Ala	Lys	Thr	Leu	Leu	Pro	Thr
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Ser	Ala	Leu	Ile	Thr	Gly	Val	Pro	Ile	Leu	Glu	Ser	Val	Gly	Ile	Ala
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Lys	Ala	Leu	Ala	Ala	Lys	Asn	Arg	Tyr	Glu	Leu	Asp	Ser	Asn	Ser	Asp
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Tyr	Pro	Ala	Thr	Gly	Ser	Phe	Ser	Arg	Ser	Ala	Val	Asn	Asn	Glu	Ser
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Glu	Ala	Lys	Thr	Gly	Leu	Ser	Gly	Leu	Ile	Thr	Gly	Ile	Ile	Ile	Gly
				405					410					415	
Cys	Ser	Leu	Leu	Phe	Leu	Thr	Pro	Met	Phe	Lys	Tyr	Ile	Pro	Gln	Cys
			420					425					430		
Ala	Leu	Ala	Ala	Ile	Val	Ile	Ser	Ala	Val	Ser	Gly	Leu	Val	Asp	Tyr
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Asp	Glu	Ala	Ile	Phe	Leu	Trp	Arg	Val	Asp	Lys	Arg	Asp	Phe	Ser	Leu
	450					455					460				
Trp	Thr	Ile	Thr	Ser	Thr	Ile	Thr	Leu	Phe	Phe	Gly	Ile	Glu	Ile	Gly
465					470					475					480
Val	Leu	Val	Gly	Val	Gly	Phe	Ser	Leu	Ala	Phe	Val	Ile	His	Glu	Ser
				485					490					495	
Ala	Asn	Pro	His	Ile	Ala	Val	Leu	Gly	Arg	Leu	Pro	Gly	Thr	Thr	Val
			500					505					510		
Tyr	Arg	Asn	Ile	Lys	Gln	Tyr	Pro	Glu	Ala	Tyr	Thr	Tyr	Asn	Gly	Ile
		515					520					525			
Val	Ile	Val	Arg	Ile	Asp	Ser	Pro	Ile	Tyr	Phe	Ala	Asn	Ile	Ser	Tyr
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Asn	Arg	Gly	Leu	Glu	Val	Asp	Arg	Ile	Asn	Phe	Val	Ile	Leu	Glu	Met
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Ser	Pro	Val	Thr	His	Ile	Asp	Ser	Ser	Ala	Val	Glu	Ala	Leu	Lys	Glu
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Leu	Tyr	Gln	Glu	Tyr	Lys	Thr	Arg	Asp	Ile	Gln	Leu	Ala	Ile	Ser	Asn

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Pro	Asn	Lys	Asp	Val	His	Leu	Thr	Ile	Ala	Arg	Ser	Gly	Met	Val	Glu
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Leu	Val	Gly	Lys	Glu	Trp	Phe	Phe	Val	Arg	Val	His	Asp	Ala	Val	Gln
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Val	Cys	Leu	Gln	Tyr	Val	Gln	Ser	Ser	Asn	Leu	Glu	Asp	Lys	His	Leu
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Ser	Phe	Thr	Arg	Arg	Tyr	Gly	Gly	Ser	Asn	Asn	Asn	Ser	Ser	Ser	Ser
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675					680					685					